

APPENDIX A

The following parameters are utilized by the AMB algorithm:

Input: X - the given design matrix (continuous + categorical) (dimension: m x n, m = # of records, n = # of predictors);
y – the dependent/target variable vector (dimension: m x 1)
Output: s – the solution vector (the model parameter vector, including the “bias” term) (dimension: (n1+1) x 1)

Step 0

For each continuous predictor

If (there is any missing observation value)
 Perform Missing Value Substitution
End

Step 1

For each continuous predictor

If (exponentially distributed)
 Log-scale the predictor and flag it
End

Detect outliers

End

Step2

// Perform Univariate Analysis for all n predictors

If (size(continuous) > 0)
 For each continuous predictor
 Calculate its Pearson’s r value (with the target)
 End
End

If (size(categorical) > 0)
 Bin the continuous target variable
 Calculate its Cramer’s V value (on the binned target groups)

End

Sort continuous predictors in Pearson’s R value

Sort categorical predictors in Cramer’s V value

// Assume n = n_conti + n_cate, n_conti = # of continuous, n_cate = # of categorical
If n_conti > 200

 Retain top 135 + ((n_conti – 200)*0.3) (30% continuous with large R values)

Else if 100 < n_conti <= 200

 Retain top 85 + ((n_conti – 100)*0.5) (50% continuous with large R values)

Else if 50 < n_conti <=100

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    Retain top 50 + ((n_conti - 50)*0.7) (70% continuous with large R values)
Else // n_conti <=50
    Retain all predictors
End
If n_cate > 200
    Retain top 135 + ((n_cate - 200)*0.3) (30% categorical with large V values)
Else if 100 < n_cate <= 200
    Retain top 85 + ((n_cate - 100)*0.5) (50% categorical with large V values)
Else if 50 < n_cate <=100
    Retain top 50 + ((n_cate - 50)*0.7) (70% categorical with large V values)
Else // n_cate <=50
    Retain all predictors
End

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Step 3

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If (size(categorical) > 0 & size(continuous)>0)
    // Merge categorical with continuous (in favor of continuous)
    Categorize continuous predictors
        For each categorical predictor c1
        For each continuous predictor c2
            Compute the Cramer's V value between c1 and c2
            If Cramer V(c1, c2) > 0.5
                Remove c1 from the retained list
            End
        End
    End
End

If (size(categorical) > 0)
    Expand all retained categorical predictors into dummies
End
If (size(categorical) > 0 && size(continuous) > 0)
    Formulate the new design matrix X by combining retained categorical and continuous
    predictors
End

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Step 4

Normalize (not z-scaling) all retained predictors (X) and obtain the new design matrix X'

Step 5

Formulate the normal equation $N = X'^T X'$ (matrix-matrix multiplication, dimension of N : n1 x n1)

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//Filter out strongly collinear predictors
While there is an off-diagonal-element of lower_triangle(X'^T X') with its absolute value > 0.8
    // assume the index is (i, j) and i > j
    Compute the correlation r_i between the target and the ith predictor

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Compute the correlation r_j between the target and the j th predictor
 If $r_i > r_j$
 Remove j th predictor from the retaining predictor list
 Else
 Remove i th predictor from the retaining predictor list
 End
 End
 If any predictor deletion (above) performed
 Reformulate the design matrix X' and the corresponding normal equation $N = X'^T X'$,
 (matrix-matrix multiplication)
 $[m, n1] = \text{size}(X')$
 End

Step 6

Perform PCA on N via SVD(N) and obtain the loading matrix M (dimension: $n1 \times n1$) and the latent vector l (dimension: $n1 \times 1$)

Step 7

If PCA successful (i.e., the SVD in PCA does not fail)

Sort the latent vector l in increasing order and obtain the sorting index;
 Use singular values l and the sorting index to identify a few bottom components C (i.e., the last d columns of M , dimension: $n1 \times d$) that represents 10 % of variance accounted for;

If ($n1 - d < 10$)

 Reformulate C by including only the last $d2 (= n1 - 10)$ columns of M

 Reset $d = d2$

End

Scan all columns/components in C and delete $d1 (<=d)$ components that don't have a predictive strength, i.e., $|\text{Pearson's } R(\text{target, component})| < 0.3$

Step 8

$k = n1 - d1$

Formulate the Mapping matrix M' from M (by removing those $d1$ components, dimension of M' : $n1 \times k$)

While ($k >= m$)

 Delete the bottom components according to the singular value

End While

Reset k to the size of remaining components

Compute $A' = X'M'$ (matrix-matrix multiplication, dimension of A' : $m \times k$)

Step 9

Append the "bias" column (all 1's) to A' as its (new) first column (dimension of A' : $m \times (k+1)$)

Pass A' to Engine (SVD + possibly a random initial guess and CGD) for component regression and generate a solution vector w (dimension: $(k+1) \times 1$)

Step 10

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// Map w back to the predictor space
  -- Compute the solution vector s = M' * w [2..k+1] (multiplication of matrix M' and a
  partial vector of w (from w[2] to w[k+1]) (dimension of s : n1 x 1)
  -- Add the "bias" term (i.e., w[1]) to s as its (new) first entry (dimension of s : (n1+1) x
  1)
Else // PCA failed
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  Append the "bias" column to X' as its (new) first column (dimension of X' : m x (n1+1))
  While (n+1 >= m)
    Delete the remaining least correlated (with target) variable
  End While
  Reset n+1 to the size of retained design matrix
  Pass all retained predictors X' to Engine (SVD + possibly a random initial guess and
CGD) for predictor regression and generate a solution vector s (dimension: (n1+1) x 1)
  End
```